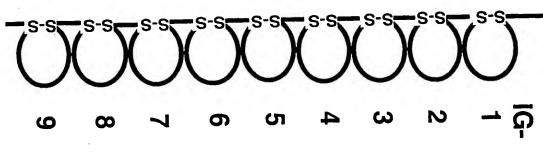


## 1 MWILALSLEGSFANVESEDLHSS

Signal Peptide

- 24 LYFVNASLQEVVFASTTGTLVPCPAAGIPPVTLRWYLATGEEIYDVPGIRHVHPNGTLQIFPFPPSSF STLIHDNTYYCTAENPSGKIRSQDVHIKAVLREPY
- 127  ${\tt TVRVEDQKTMRGNVAVFK} \textbf{C} {\tt IPSSVEAYITVVSWEKDTVSLVSGSRFLITSTGALYIKDVQNEDGLYN$ YR**C**ITRHRYTGETRQSNSARLFVSDPANSAP
- 226 SILDGFDHRKAMAGQRVELPCKALGHPEPDYRWLKDNMPLELSGRFQKTVTGLLIENIRPSDSGSYVC EVSNRYGTAKVIGRLYVKQPLKA
- 317 TISPRKVKSSVGSQVSLSCSVTGTEDQELSWYRNGEILNPGKNVRITGINHENLIMDHMVKSDGGAYQ CFVRKDKLSAQDYVQVVLEDGTPKI
- 410 ISAFSEKVVSPAEPVSLMCNVKGTPLPTITWTLDDDPILKGGSHRISQMITSEGNVVSYLNISSSQVR DGGVYR**C**TANNSAGVVLYQARINVRGPAS
- 507  ${\tt IRPMKNITAIAGRDTYIH}{\tt CRVIGYPYYSIKWYKNSNLLPFNHRQVAFENNGTLKLSDVQKEVDEGEYT}$ CNVLVQPQLSTSQSVHVTVKVPPFIQPFE
- 604 FPRESIGQRVFIPCVVVSGDLPITITWQKDGRPIPGSLGVTIDNIDFTSSLRISNLSLMHNGNYTCIA RNEAAAVEHQSQLIVRVPPKFVVQPR
- 698 DQDGIYGKAVILNCSAEGYPVPTIVWKFSKGAGVPQFQPIALNGRIQVLSNGSLLIKHVVEEDSGYYL CKVSNDVGADVSKSMYLTVKIPAMITS
- 793 YPNTTLATQGQKKEMSCTAHGEKPIIVRWEKEDRIINPEMARYLVSTKEVGEEVISTLQILPTVREDS GFFSCHAINSYGEDRGIIQLTVQEPPD

## FIG. 2A





888 PPEIEIKDVKARTITLRWTMGFDGNSPITGYDIECKNKSDSWDSAQRTKDVSPQLNSATIIDIHPSST YSIRMYAKNRIGKSEPSNELTITADEAA

984 PDGPPQEVHLEPISSQSIRVTWKAPKKHLQNGIIRGYQIGYREYSTGGNFQFNIISVDTSGDSEVYTL DNLNKFTQYGLVVQACNRAGTGPSSQEIITTTLED

1087 VPSYPPENVQAIATSPESISISWSTLSKEALNGILQGFRVIYWANLMDGELGEIKNITTTQPSLELDG LEKYTNYSIQVLAFTRAGDGVRSEQIFTRTK

1186 EDVPGPPAGVKAAAASASMVFVSWLPPLKLNGIIRKYTVFCSHPYPTVISEFEASPDSFSYRIP**NLS**R NRQYSVWVVAVTSAGRGNSSEIITVEPL

1282 AKAPARILTESGTVTTPWMKDIVLPCKAVGDPSPAVKWMKDSNGTPSLVTIDGRRSIFSNGSFIIRTV KAEDSGYYSCIANNNWGSDEIILNLQ

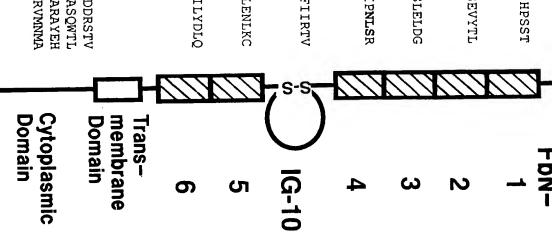
1376 VQVPPDQPRLTVSKTTSSSITLSWLPGDNGGSSIRGYILQYSEDNSEQWGSFPISPSERSYRLENLKC **GTWYKFTLTAQNGVGPGRISEIIEAKTL** 

1472 GKEPQFSKEQELFASINTTRVRLNLIGWNDGGCPITSFTLEYRPFGTTVWTTAQRTSLSKSYILYDLQ EATWYELQMRVCNSAGCAEKQANFATLNYDGSTIPPLIKSVVQNEEGLTTNEGLK

1595 MLVTISCILVGVLLLEVLLLVV

1617 RRRREQRLKRLRDAKSLAEMLMSKNTRTSDTLSKQQQTLRMHIDIPRAQLLIEERDTMETIDDRSTV AKMEEQLRHAKFTITECFISDTSSEQLTAGTNEYTDSLTSSTPSESGICRFTASPPKPQDGGRVMNMA NRPHPTISAHTLTTDWRLPTPRAAGSVDKESDSYSVSPSQDTDRARSSMVSTESASSTYEELARAYEH VPKAIGQVTSYICLHTLEWTFC LLTDADFGEAAKQKSLTVTHTVHYQSVSQATGPLVDVSDARPGTNPTTRRNAKAGPTARNRYASQWTL

FIG. 2B





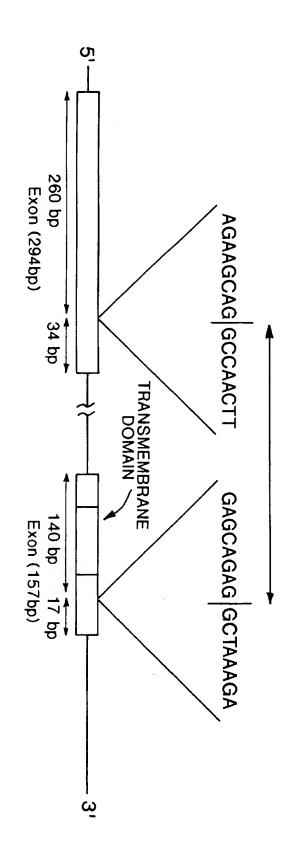
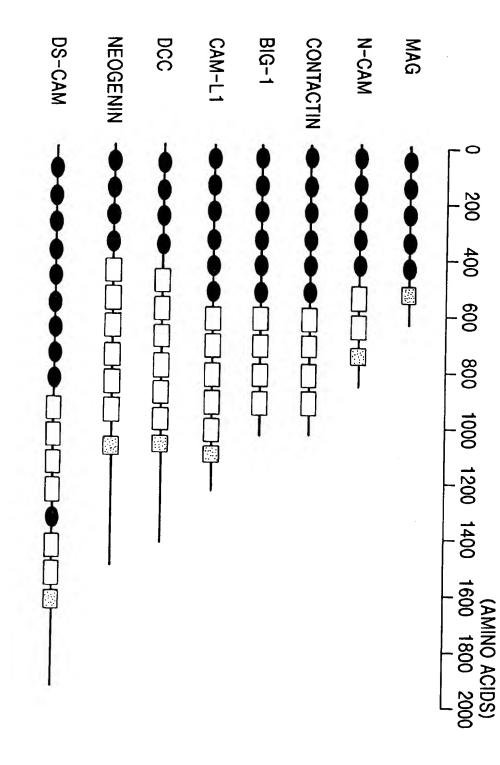


FIG. 3





Ig TYPE-C2 DOMAIN
TRANSMEMBRANE DOMAIN

FIBRONECTIN TYPE-III DOMAIN

FIG. 4